

### **INPUT SET: S23914.raw**

**This Raw Listing contains the General Information Section and up to the first 5 pages.**

## SEQUENCE LISTING

(1) General Information

(i) APPLICANT: Jung, Rudolf  
Beach, Larry R.  
Dress, Virginia M.  
Rao, A. Gururaj  
Ranch, Jerome P.  
Ertl, David S.  
Higgins, Regina K.

--  
13 (ii) TITLE OF THE INVENTION: Alteration of Amino Acid Compositions  
14 in Seeds

(iii) NUMBER OF SEQUENCES: 13

(iv) CORRESPONDENCE ADDRESS:

(A) ADDRESSEE: Pioneer Hi-Bred International, Inc.  
(B) STREET: 7100 NW 62nd Avenue, P.O. Box 1000  
(C) CITY: Johnston  
(D) STATE: IA  
(E) COUNTRY: USA  
(F) ZIP: 50131

(v) COMPUTER READABLE FORM:

- (A) MEDIUM TYPE: Diskette
- (B) COMPUTER: IBM Compatible
- (C) OPERATING SYSTEM: DOS
- (D) SOFTWARE: FastSEQ for Windows Version 2.0

(vi) CURRENT APPLICATION DATA:

- (A) APPLICATION NUMBER:
- (B) FILING DATE:
- (C) CLASSIFICATION:

(vii) PRIOR APPLICATION DATA:

(A) APPLICATION NUMBER:  
(B) FILING DATE:

(viii) ATTORNEY/AGENT INFORMATION:

(A) NAME: Michel, Marianne H  
(B) REGISTRATION NUMBER: 35,286  
(C) REFERENCE/DOCKET NUMBER: 0815

RAW SEQUENCE LISTING  
PATENT APPLICATION US/09/020,716DATE: 02/27/98  
TIME: 18:53:29

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47  
 48       (ix) TELECOMMUNICATION INFORMATION:  
 49        (A) TELEPHONE: 515-334-4467  
 50        (B) TELEFAX: 515-334-6883  
 51        (C) TELEX:  
 52  
 53  
 54        (2) INFORMATION FOR SEQ ID NO:1:  
 55  
 56        (i) SEQUENCE CHARACTERISTICS:  
 57        (A) LENGTH: 3363 base pairs  
 58        (B) TYPE: nucleic acid  
 59        (C) STRANDEDNESS: single  
 60        (D) TOPOLOGY: linear  
 61  
 62        (ii) MOLECULE TYPE: Other  
 63  
 64        (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1:  
 65  
 66 TCGACCTCGA GGGGGGGCCC GGTACCCAGC TTTTGTTCCTC TTAGTGAGG GTTAATTGCG 60  
 67 CGCTTGGCGT AATCATGGTC ATAGCTGTTT CCTGTGTGAA ATTGTTATCC GCTCACAAATT 120  
 68 CCACACAAACA TACGACCCGG AACCATAAAG TGAAAGCCT GGGGTGCCTA ATGAGTGAGC 180  
 69 TAACTCACAT TAATTGCGTT GCGCTCACTG CCCGCTTCC AGTCGGGAAA CCTGTCGTGC 240  
 70 CAGCTGCATT AATGAATCGG CCAACGCGG GGGAGAGGCG GTTGTGCTAT TGGGCGCTCT 300  
 71 TCCGCTTCCT CGCTCACTGA CTCGCTGCCTC TCGGTCGTTC GGCTGCGGCG AGCGGTATCA 360  
 72 GCTCACTCAA AGGGGTAAT ACGGTTATCC ACAGAACATCAG CGGATAACGC AGGAAAGAAC 420  
 73 ATGTGAGCAA AAGGCCAGCA AAAGGCCAGG AACCGTAAAA AGGCCGCGTT GCTGGCGTTT 480  
 74 TTCCATAGGC TCCGCCCCCC TGACGAGCAT CACAAAAAATC GACGCTCAAG TCAGAGGTGG 540  
 75 CGAAACCCGA CAGGACTATA AAGATACCAAG GCGTTTCCCC CTGGAAGCTC CCTCGTGC 600  
 76 TCTCTGTTC CGACCCCTGCC GCTTACCGGA TACCTGTCCC CCTTCTCTCCC TTGGGAAGC 660  
 77 GTGGCGCTTT CTCATAGCTC ACCTGTAGG TATCTCAGTT CGGTGTAGGT CGTTCGCTCC 720  
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 79 TATCGTCTTG AGTCCAACCC GGTAAAGACAC GACTTATCGC CACTGGCAGC AGCCACTGGT 840  
 80 AACAGGATTA GCAGAGCGAG GTATGTAGGC GGTGCTACAG AGTTCTTGAA GTGGTGGCCT 900  
 81 AACTACGGCT ACACTAGAAG GACAGTATTG GGTATCTCGC CTCTGCTGAA GCCAGTTACC 960  
 82 TTGGGAAAAA GAGTTGGTAG CTCTTGATCC GGCAAAACAAA CCACCGCTGG TAGCGGTGGT 1020  
 83 TTTTTGTTT GCAAGCAGCA GATTACGCG AGAAAAAAAAG GATCTCAAGA AGATCCTTTG 1080  
 84 ATCTTTCTA CGGGGTCTGA CGCTCAGTGG AACGAAAAC CACGTTAAGG GATTTTGGTC 1140  
 85 ATGAGATTAT CAAAAAGGAT CTCACCTAG ATCCTTTAA ATTAAAAATG AAGTTTTAAA 1200  
 86 TCAATCTAAA GTATATATGA GTAAACTTGG TCTGACAGTT ACCAATGCTT AATCAGTGAG 1260  
 87 GCACCTATCT CAGCGATCTG TCTATTCGT TCATCCATAG TTGCTGACT CCCCGTCGTG 1320  
 88 TAGATAACTA CGATACGGGA GGGCTTACCA TCTGGCCCCA GTGCTGCAAT GATACCGCGA 1380  
 89 GACCCACGCT CACCGGTCTCC AGATTATCA GCAATAAACCC AGCCAGCCGG AAGGGCCGAG 1440  
 90 CGCAGAAGTG GTCCTGCAAC TTTATCCGCC TCCATCCAGT CTATTAATTG TTGCGGGAA 1500  
 91 GCTAGAGTAA GTAGTTGCC AGTTAATAGT TTGCGCAACG TTGTTGCCAT TGCTACAGGC 1560  
 92 ATCGTGGTGT CACGCTCGTC GTTTGGTATG GCTTCATTCA GCTCCGGTTC CCAACGATCA 1620  
 93 AGGCAGTTA CATGATCCCC CATGTTGTGC AAAAAAGCGG TTAGCTCCTT CGGTCTCCG 1680  
 94 ATCGTTGTCA GAAGTAAGTT GGGCGCAGTG TTATCACTCA TGTTATGGC AGCACTGCAT 1740  
 95 AATTCTCTTA CTGTATGCC ATCCGTAAGA TGCTTTCTG TGACTGGTGA GTACTCAACC 1800  
 96 AAGTCATTCT GAGAATAGTG TATGCGGCCA CCGAGTTGCT CTTGCCGGC GTCAATACGG 1860  
 97 GATAATACCG CGCCACATAG CAGAACTTTA AAAGTGCTCA TCATTGGAAA ACGTTCTTCG 1920  
 98 GGGCGAAAAC TCTCAAGGAT CTTACCGCTG TTGAGATCCA GTTCGATGTA ACCCACTCGT 1980  
 99 GCACCCAACCT GATCTTCAGC ATCTTTTACT TTCACCAGCG TTTCTGGTG AGCAAAACAA 2040

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PATENT APPLICATION US/09/020,716DATE: 02/27/98  
TIME: 18:53:32

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102	ATATTGAAAT	GTATTTAGAA	AAATAAACAA	ATAGGGTTC	CGCGCACATT	TCCCCGAAAA	2220
103	GTGCCACCTA	AATTGTAAGC	GTAAATATTT	TGTTAAAATT	CGCGTTAAAT	TTTTGTTAAA	2280
104	TCAGCTCATT	TTTTAACCAA	TAGGCCGAAA	TCGGCAAAAT	CCCTTATAAA	TCAAAGAACAT	2340
105	AGACCGAGAT	AGGGTTGAGT	GTTGTTCCAG	TTTGGAACAA	GAGTCCACTA	TTAAAGAACG	2400
106	TGGACTCCAA	CGTCAAAGGG	CGAAAAACCG	TCTATCAGGG	CGATGGCCCA	CTACGTGAAC	2460
107	CATCACCCCTA	ATCAAGTTTT	TTGGGGTCGA	GGTGCCTAA	AGCACTAAAT	CGGAACCCCTA	2520
108	AAGGGAGCCC	CCGATTAGA	GCTTGACGGG	GAAAGCCGC	GAACGTGGCG	AGAAAGGAAG	2580
109	GGAAGAACG	GAAAGGAGCG	GGCGCTAGGG	CGCTGGCAAG	TGTAGCGGTC	ACGCTGGCG	2640
110	TAACCACCAAC	ACCCGCCGCG	CTTAATGCGC	CGCTACAGGG	CGCGTCCCAT	TCGCCATTCA	2700
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113	GACGTTGTAA	AACGACGGCC	AGTGAGCGCG	CGTAATACGA	CTCACTATAG	GGCGAATTGG	2880
114	AGCTCCACCG	CGGTGGCGGC	CGCTCTAGAA	CTAGTGGATC	CGTCGACTAG	AGGGCCCGAC	2940
115	GTCGAACCTTA	GGCACTAAGG	GATGTGAGGC	CAGCATCACC	GTTGCAGAAA	TTGACACAAG	3000
116	CATCACCAACA	ATTTTCCAAA	TAGAGTTTC	TTTCTTCGTC	GTCAGCAGCT	GCGTTGACCA	3060
117	TGTAGTCACA	CATGGAAGCC	CTACACCCCA	AGTTGCAATA	CTTGACGGTG	TCTGGTTCAT	3120
118	CTGAGTTGGA	CACAAGGGCC	AATTGGGGA	AGCCTGTAGG	GCATTTCCG	CTACTTGTGA	3180
119	GTTTACACCT	ACAGACCCCT	GGCATAACT	TCTGAGCACC	ACGGACGCAG	CAAAGGTTGT	3240
120	AGCAGTTCT	TCCTAGGGT	CTCCTGCAGC	AACTCTTGCC	TTCTACTTGC	ACCTGTTCGA	3300
121	GAACCAACCC	CAGTATAAGT	AAACACACCA	TCACACCCCT	GAGGCCCTTG	CTGGTGGCCA	3360
122	TGG						3363
123							

## 124 (2) INFORMATION FOR SEQ ID NO:2:

125

## 126 (i) SEQUENCE CHARACTERISTICS:

127

- (A) LENGTH: 3365 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

131

## 132 (ii) MOLECULE TYPE: Other

133

## 134 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:2:

135

136	TCGACCTCGA	GGGGGGGCC	GGTACCCAGC	TTTGTTCCC	TTTAGTGAGG	GTAAATTGCG	60
137	CGCTTGGCGT	AATCATGGTC	ATAGCTGTT	CCTGTGTGAA	ATTGTTATCC	GCTCACAAATT	120
138	CCACACAACA	TACGAGCCGG	AACCATAAAG	TGTAAAGCCT	GGGGTGCCTA	ATGAGTGAGC	180
139	TAACTCACAT	TAATTGCGTT	GGCCTCACTG	CCCGCTTCC	AGTCGGAAA	CCTGTCGTGC	240
140	CAGCTGCATT	AATGAATCGG	CCAACGCGCG	GGGAGAGGCG	GTTGCGTAT	TGGCGCTCT	300
141	TCCGCTTCCT	CGCTCACTGA	CTCGCTGC	TCGGTCGTT	GGCTGCGCG	AGCGGTATCA	360
142	GCTCACTCAA	AGGCGGTAAAT	ACGGTTATCC	ACAGAACATAG	GGGATAACGC	AGGAAAGAAC	420
143	ATGTGAGCAA	AAGGCCAGCA	AAAGGCCAGG	AACCGTAAAAA	AGGCCGCGTT	GCTGGCGTT	480
144	TTCCATAGGC	TCCGCCCCCC	TGACGAGCAT	CACAAAAATC	GACGCTCAAG	TCAGAGGTGG	540
145	CGAAACCCGA	CAGGACTATA	AAGATACCAAG	GCGTTTCCCC	CTGGAAGCTC	CCTCGTGC	600
146	TCTCCTGTT	CGACCCCTGCC	GCTTACCGGA	TACCTGTCCG	CCTTCCTCCC	TTCGGGAAGC	660
147	GTGGCGCTTT	CTCATAGCTC	ACGCTGTAGG	TATCTCAGTT	CGGTGTAGGT	CGTTCGCTCC	720
148	AAGCTGGCT	GTGTGCAACGA	ACCCCCCGTT	CAGCCCGACC	GCTGCGCCTT	ATCCGTAAC	780
149	TATCGTCTTG	AGTCCAACCC	GGTAAGACAC	GACTTATCGC	CACTGGCAGC	AGCCACTGGT	840
150	AACAGGATTA	GCAGAGCGAG	GTATGTAGGC	GGTGCTACAG	AGTTCTTGAA	GTGGTGGCCT	900
151	AACTACGGCT	ACACTAGAAG	GACAGTATT	GGTATCTGCG	CTCTGCTGAA	GCCAGTTACC	960
152	TTCGGAAAAAA	GAGTTGGTAG	CTCTTGATCC	GGCAAACAAA	CCACCGCTGG	TAGCGGTGGT	1020

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PATENT APPLICATION US/09/020,716DATE: 02/27/98  
TIME: 18:53:35

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155	ATGAGATTAT CAAAAAGGAT CTTCACCTAG ATCCTTTAA ATTAAAATG AAGTTTAAA	1200
156	TCAATCTAAA GTATATATGA GTAAACTTGG TCTGACAGTT ACCAATGCTT AATCAGTGAG	1260
157	GCACCTATCT CAGCGATCTG TCTATTTCTG TCATCCATAG TTGCCTGACT CCCCGTCGTG	1320
158	TAGATAACTA CGATACGGGA GGGCTTACCA TCTGGCCCA GTGCTGCAAT GATACCGCGA	1380
159	GACCCACGCT CACCGGCTCC AGATTTATCA GCAATAAACCC AGCCAGCCGG AAGGGCCGAG	1440
160	CGCAGAAAGTG GTCCTGCAAC TTTATCCGCC TCCATCCAGT CTATTAATTG TTGCGGGAA	1500
161	GCTAGAGTAA GTAGTCGCC AGTTAATAGT TTGCGCAACG TTGTTGCCAT TGCTACAGGC	1560
162	ATCGTGGTGT CACGCTCGTC GTTTGGTATG GCTTCATTCA GCTCCGGTTC CCAACGATCA	1620
163	AGCGAGTTA CATGATCCCC CATGTTGTGC AAAAAAGCGG TTAGCTCCTT CGGTCTCCG	1680
164	ATCGTGTCA GAAGTAAGTT GGGCGCAGTG TTATCACTCA TGGTTATGGC AGCACTGCAT	1740
165	AATTCTCTTA CTGTCATGCC ATCCGTAAGA TGCTTTCTG TGACTGGTGA GTACTCAACC	1800
166	AAGTCATTCT GAGAATAGTG TATGCGGCGA CCGAGTTGCT CTTGCGCCGGC GTCAATACGG	1860
167	GATAATACCG CGCCACATAG CAGAACTTTA AAAGTGCCTCA TCATTGGAAA ACGTTCTTCG	1920
168	GGCGAAAAC TCTCAAGGAT CTTACCGCTG TTGAGATCCA GTTCGATGTA ACCCACTCGT	1980
169	GCACCCAACG GATCTTCAGC ATCTTTACT TTCACCCAGCG TTTCTGGGTG AGCAAAAACA	2040
170	GGAAGGCAAA ATGCCGCAAA AAAGGGAATA AGGGCGACAC GGAAATGTTG AATAACTCATA	2100
171	CTCTTCTTT TTCAATATTA TTGAAGCATT TATCAGGGTT ATTGTCTCAT GAGCGGATAAC	2160
172	ATATTTGAAT GTATTTAGAA AAATAAACAA ATAGGGGTT CGCGCACATT TCCCCGAAAA	2220
173	GTGCCACCTA AATTGTAAGC GTTAATATTT TGTAAAATT CGCGTTAAAT TTTTGTAAA	2280
174	TCAGCTCATT TTTAACCAA TAGGCGAAA TCGGCAAAAT CCCTTATAAAA TCAAAAGAAT	2340
175	AGACCGAGAT AGGGTTGAGT GTTGTCCAG TTTGGAACAA GAGTCCACTA TTAAAGAACG	2400
176	TGGACTCCAA CGTCAAAGGG CGAAAAACCG TCTATCAGGG CGATGGCCCA CTACGTGAAC	2460
177	CATCACCCCTA ATCAAGTTT TTGGGTCGA GGTGCCGTAA AGCACTAAAT CGGAACCCCTA	2520
178	AAGGGAGCCC CCGATTAGA GCTTGACGGG GAAAGCCGGC GAACGTGGCG AGAAAGGAAG	2580
179	GGAAGAAAGC GAAAGGAGCG GGCCTAGGG CGCTGGCAAG TGTAGCGGTG ACGCTGCGCG	2640
180	TAACCACCAC ACCCGCCCGC CTTAATGCGC CGCTACAGGG CGCGTCCCAT TCGCCATTCA	2700
181	GGCTGCGCAA CTGTTGGAA GGGCGATCGG TGCGGGCCTC TTCGCTATTAA CGCCAGCTGG	2760
182	CGAAAGGGGG ATGTGCTGCA AGGCATTAA GTTGGGTAAC GCCAGGGTT TCCCAGTCAC	2820
183	GACGTTGTAA AACGACGGCC AGTGAGCGCG CGTAATACGA CTCACTATAG GCGAATTGG	2880
184	AGCTCCACCG CGGTGGCGC CGCTCTAGAA CTAGTGGATC CGTCGACTAG AGGGCCCGAC	2940
185	GTCGAACCTTA GGCACTAAGG GATGTGAGGC CAGCATCACC GTTGCAGAAA TTGACACAAG	3000
186	CATCACCCACA ATTTTCCAAA TAGAGTTCA TTTCTCGTC GTCAGCAGCT GCGTTGACCA	3060
187	TGTAGTCACA CATGGAAGCC CTACACCCCA AGTTGCAATA CTTGACGGTG TCTGGTTCAT	3120
188	CTGAGTTGGA CACAAGGGCC AATTGGGAA AGCCTTTCGG GCATTTTCCG CTACTAGTCA	3180
189	GCTTACACTT GCAGACGCCCT GCGCAAAGCT TCTTGGCGCC TTGACTTTG CAAAGGTTGT	3240
190	AGCACTTCCT TCCCAGGGTA CTCTTGCAGC AACTCTTGCC TTCTACTTGC ACCTGTTCGA	3300
191	GAACCAACCC CAGTATAAGT AAACACACCA TCACACCCCTT GAGGCCCTTG CTGGTGGCCA	3360
192	TGGTG	3365

193  
194 (2) INFORMATION FOR SEQ ID NO:3:195  
196 (i) SEQUENCE CHARACTERISTICS:  
197 (A) LENGTH: 5360 base pairs  
198 (B) TYPE: nucleic acid  
199 (C) STRANDEDNESS: single  
200 (D) TOPOLOGY: linear201  
202 (ii) MOLECULE TYPE: Other203  
204 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:3:

205

RAW SEQUENCE LISTING  
PATENT APPLICATION US/09/020,716DATE: 02/27/98  
TIME: 18:53:39

## INPUT SET: S23914.raw

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207	ATTTTTAAC	CAATAGGCCG	AAATCGCAA	AATCCCTTAT	AAATCAAAAG	AATAGACCGA	120
208	GATAGGGTTG	AGTGGTGTTC	CAGTTGGAA	CAAGAGTCCA	CTATTAAGA	ACGTGGACTC	180
209	CAACGTCAAA	GGGCGAAAAA	CCGTCTATCA	GGGCGATGGC	CCACTACGTG	AACCATCACC	240
210	CTAATCAAGT	TTTTTGGGGT	CGAGGTGCCG	TAAAGCACTA	AATCGGAACC	CTAAAGGGAG	300
211	CCCCCGATT	AGAGCTTGAC	GGGGAAAGCC	GGCGAACGTG	GCGAGAAAGG	AAGGGAAGAA	360
212	AGCGAAAGGA	GCGGGCGCTA	GGCGCCTGGC	AAAGTGTAGCG	GTCACGCTGC	GCGTAACCAC	420
213	CACACCGCC	GCGCTTAATG	CGCCGCTACA	GGGCGCGTCC	CATTGCCAT	TCAGGCTGCG	480
214	CAACTGTTGG	GAAGGGCGAT	CGGTGCGGGC	CTCTTCGCTA	TTACGCCAGC	TGGCGAAAGG	540
215	GGGATGTGCT	GCAAGGGCGAT	TAAGTTGGGT	AACGCCAGGG	TTTCCCAGT	CACGACGTTG	600
216	TAAAACGACG	GCCAGTGAGC	GCCGTAATA	CGACTCACTA	TAGGGCGAAT	TGGAGCTCCA	660
217	CCGGGGTGGC	GGCCGCTCTA	GATTATATAA	TTTATAAGCT	AAACAACCCG	GCCCTAAAGC	720
218	ACTATCGTAT	CACCTATCTA	AATAAGTCAC	GGGAGTTTCG	AACGTCCACT	TCGTCGCACG	780
219	GAATTGCATG	TTCTTGTGTT	GAAGCATATT	CACGCAATCT	CCACACATAA	AGGTTTATGT	840
220	ATAAAACTTAC	ATTTAGCTCA	GTTTAATTAC	AGTCTTATTT	GGATGCATAT	GTATGTTCT	900
221	CAATCCATAT	AAGTTAGAGT	AAAAAAATAAG	TTTAAATTTT	ATCTTAATTC	ACTCCAACAT	960
222	ATATGGATCT	ACAATACTCA	TGTGCATCCA	AACAAACTAC	TTATATTGAG	GTGAATTGAG	1020
223	TAGAAAATTAA	ACTAACTTAC	ACACTAAGCC	AATCTTTACT	ATATTAAAGC	ACCAGTTCA	1080
224	ACGATCGTCC	CGCGTCAATA	TTATTAAAAA	ACTCCTACAT	TTCTTTATAA	TCAACCCGCA	1140
225	CTCTTATAAT	CTCTTCTCTA	CTACTATAAT	AAGAGAGTTT	ATGTACAAAAA	TAAGGTGAAA	1200
226	TTATCTATAA	GTGTTCTGG	TATTGGTTGT	TGGCTCCCAT	ATTCACACAA	CCTAATCAAT	1260
227	AGAAAACATA	TGTTTTATT	AAACAAAATT	TATCATATAT	CATATATATA	TATATATCAT	1320
228	ATATATATAT	AAACCGTAGC	AATGCACGGG	CATATAACTA	GTGCAACTTA	ATACATGTGT	1380
229	GTATTAAGAT	GAATAAGAGG	GTATCCAAT	AAAAAAACTTG	TTGCTTACGT	ATGGATCGAA	1440
230	AGGGGTTGG	AACGATTAAA	CGATTAAATC	TCTTCCTAGT	AAAATTGAA	TAGAAGGAGA	1500
231	TTTAATATAT	CCCAATCCCC	TTCGATCATC	CAGGTGCAAC	CGTATAAGTC	CTAAAGTGGT	1560
232	GAGGAACACG	AAAGAACCAT	GCATTGGCAT	GTAAAGCTCC	AAGAATTGTT	TGTATCCTTA	1620
233	ACAACTCACA	GAACATCAAC	CAAAATTGCA	CGTCAAGGGT	ATTGGGTAAG	AAACAATCAA	1680
234	ACAAATCCTC	TCTGTGTGCA	AAGAAACACG	GTGAGTCATG	CCGAGATCAT	ACTCATCTGA	1740
235	TATACATGCT	TACAGCTCAC	AAAGACATTAC	AAACAACCTA	TATTGCATTA	CAAAGATCGT	1800
236	TTCATGAAAA	ATAAAATAGG	CCGGACAGGA	AAAAAACTCT	TGACGTGTAA	AGTAAATTAA	1860
237	CAACAAAAAA	AAAGCCATAT	GTCAAGCTAA	ATCTAATTG	TTTACGTAG	ATCAACAAACC	1920
238	TGTAGAAGGC	AACAAAACGT	AGCCACGCG	AGTACAGAA	TGATTCCAGA	TGAACCATCG	1980
239	ACGTGCTACG	TAAGAGACGT	GACCGAGTCAT	ATACATTG	CAAGAAACCA	TGAAGCTGCC	2040
240	TACAGCCGTC	TCGGTGGCAT	AAGAACACAA	GAAATTGTT	TAATTAATCA	AAGCTATAAA	2100
241	TAACGCTCGC	ATGCCTGTG	ACTTCTCCAT	CACCACCACT	GGGTCTTCAG	ACCATTAGCT	2160
242	TTATCTACTC	CAGAGCCAG	AAGAACCGA	TCGACACCAT	GGCCACCAGC	AAGGGCCTCA	2220
243	AGGGTGTGAT	GGTGTGTTA	CTTATACTGG	GGTTGGTTCT	CGAACAGGTG	CAAGTAGAAC	2280
244	GCAAGAGTTG	CTGCAACAGT	ACCCCTGGAA	GGAAGTGTCA	CAACCTTTGC	AAAGTCAAAG	2340
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247	ATTGCAACTT	GGGGTGTAGG	GCTTCCATGT	GTGACTACAT	GGTCAACGCA	GCTGCTGACG	2520
248	ACGAAGAAAT	GAAACTCTAT	TTGGAAAATT	GTGGTGTATGC	TTGTGTCAAT	TTCTGCAACG	2580
249	GTGATGCTGG	CCTCACATCC	CTTAGTGCCT	AAGTTGACG	TCGGGCCCTC	TAGTCGACGG	2640
250	ATCCCCGGCG	GTGTCCCCCA	CTGAAGAAAC	TATGTGCTGT	AGTATAGCCG	CTGCCCCGCTG	2700
251	GCTAGCTAGC	TAGTTGAGTC	ATTAGCGGC	GATGATTGAG	TAATAATGTG	TCACGCATCA	2760
252	CCATGCATGG	GTGGCAGTGT	CAGTGTGAGC	AATGACCTGA	ATGAACAATT	GAAATGAAAAA	2820
253	GAAAAAAAGTA	TTGTTCCAAA	TTAAACGTTT	TAACCTTTA	ATAGGTTTAT	ACAATAATTG	2880
254	ATATATGTTT	TCTGTATATG	TCTAATTG	TATCATCCAT	TTAGATATAG	ACAAAAAA	2940
255	ATCTAAGAAC	TAAAACAAAT	GCTAATTG	AATGAAGGG	GTATATATTG	GGATAATGTC	3000
256	GATGAGATCC	CTCGTAATAT	CACCGACATC	ACACGTGTCC	AGTTAATGT	TCAGTGATAC	3060
257	GTGTATTAC	ATTTGTTGCG	CGTAGGCGTA	CCCAACAAATT	TTGATCGACT	ATCAGAAAGT	3120
258	CAACGGAAGC	GAGTCGACCT	CGAGGGGGGG	CCCGGTACCC	AGCTTTGTT	CCCTTAGTG	3180